## SEQUENCE LISTING

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<110> Perez-Villar, Juan J.
     Chang, Han
     Yang, Wen-Pin
     Wu, Yuli
     Whitney, Gena S.
      Kanner, Steven B.
<120> Identification and Cloning of a Full-length Human
      Clnk-related Gene, MIST (Mast Cell Immunoreceptor
      Signal Transducer)
<130> 3053-4113US1
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<150> 60/237030
<151> 2000-09-29
<160> 52
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cattggacag ccgacctgga acacacagac gaggttggaa agagtggaca aacccatttc 780
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<211> 443

<212> PRT

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<223> HUMAN FULL-LENGTH MIST CDNA CLONE #8, TRANSLATED AMINO ACID SEQUENCE

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Asn Ser Ala Thr Gly Gln Tyr Gln Arg Met Asn Lys Pro Leu Leu Asp
50 55 60

Trp Glu Arg Asn Phe Ala Ala Val Leu Asp Gly Ala Lys Gly His Ser
65 70 75 80

Asp Asp Asp Tyr Asp Asp Pro Glu Leu Arg Met Glu Glu Thr Trp Gln 85 90 95

Ser Ile Lys Ile Leu Pro Ala Arg Pro Ile Lys Glu Ser Glu Tyr Ala 100 105 110

Asp Thr His Tyr Phe Lys Val Ala Met Asp Thr Pro Leu Pro Leu Asp 115 120 125

Thr Arg Thr Ser Ile Ser Ile Gly Gln Pro Thr Trp Asn Thr Gln Thr 130 135 140

Arg Leu Glu Arg Val Asp Lys Pro Ile Ser Lys Asp Val Arg Ser Gln 145 150 155 160

Asn Ile Lys Gly Asp Ala Ser Val Arg Lys Asn Lys Ile Pro Leu Pro 165 170 175

Pro Pro Arg Pro Leu Ile Thr Leu Pro Lys Lys Tyr Gln Pro Leu Pro 180 185 190

Pro Glu Pro Glu Ser Ser Arg Pro Pro Leu Ser Gln Arg His Thr Phe
195 200 205

Pro Glu Val Gln Arg Met Pro Ser Gln Ile Ser Leu Arg Asp Leu Ser 210 215 220

Glu Val Leu Glu Ala Glu Lys Val Pro His Asn Gln Arg Lys Pro Glu 225 230 235 240

Ser Thr His Leu Leu Glu Asn Gln Asn Thr Gln Glu Ile Pro Leu Ala 245 250 255

Ile Ser Ser Ser Ser Phe Thr Thr Ser Asn His Ser Val Gln Asn Arg 260 265 270

Asp His Arg Gly Gly Met Gln Pro Cys Ser Pro Gln Arg Cys Gln Pro
275 280 285

Pro Ala Ser Cys Ser Pro His Glu Asn Ile Leu Pro Tyr Lys Tyr Thr 290 295 300

Ser Trp Arg Pro Pro Phe Pro Lys Arg Ser Asp Arg Lys Asp Val Gln 305 310 315 320

His Asn Glu Trp Tyr Ile Gly Glu Tyr Ser Arg Gln Ala Val Glu Glu 325 330 335

Ala Phe Met Lys Glu Asn Lys Asp Gly Ser Phe Leu Val Arg Asp Cys 340 345 350

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Ser Thr Lys Ser Lys Glu Glu Pro Tyr Val Leu Ala Val Phe Tyr Glu
355 360 365
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Asn Lys Val Tyr Asn Val Lys Ile Arg Phe Leu Glu Arg Asn Gln Gla 370 375 380

Phe Ala Leu Gly Thr Gly Leu Arg Gly Asp Glu Lys Phe Asp Ser Val 385 390 395 400

Glu Asp Ile Ile Glu His Tyr Lys Asn Phe Pro Ile Ile Leu Ile Asp 405 410 415

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Pro Leu Pro Leu Thr Arg His Leu Leu Pro Leu 435 440

<210> 3

<211> 2335

<212> DNA

<213> HUMAN

<220>

<223> HUMAN MIST SPLICE VARIANT CDNA CLONE #7, NUCLEIC ACID SEQUENCE

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tctactaaaa atacaaaaaa tcagctgggc gtggtggtgg gtgcctgtag tcccagctac 2220
tegggagget gaggeaggag aatggtgtga accegggagg eggagettge agtgageega 2280
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<210> 4

<211> 428

<212> PRT

<213> HUMAN

<220>

<223> HUMAN MIST SPLICE VARIANT CLONE #7, AMINO ACID SEQUENCE

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Leu Lys Phe Gln Asn Phe Ser Leu Pro Lys Asn Arg Ser Trp Pro Arg 25 20

Ile Asn Ser Ala Thr Gly Gln Tyr Gln Arg Met Asn Lys Pro Leu Leu 45 40 35

Asp Trp Glu Arg Asn Phe Ala Ala Val Leu Asp Gly Ala Lys Gly Fis 55

Ser Asp Asp Asp Tyr Asp Asp Pro Glu Leu Arg Met Glu Glu Thr Trp 75 70 65

Gln Ser Ile Lys Ile Leu Pro Ala Arg Pro Ile Lys Glu Ser Glu Tyr

85 90 95

Ala	Asp	Thr	His	Tyr	Phe	Lys	Val	Ala	Met	Asp	Thr	Pro	Leu	Pro	Leu
			100					105					110		

- Asp Thr Arg Thr Ser Ile Ser Ile Gly Gln Pro Thr Trp Asn Thr Gln 115 120 125
- Thr Arg Leu Glu Arg Val Asp Lys Pro Ile Ser Lys Asp Val Arg Ser 130 135 140
- Pro Pro Pro Arg Pro Leu Ile Thr Leu Pro Lys Lys Tyr Gln Pro Leu 165 170 175
- Pro Pro Glu Pro Glu Ser Ser Arg Pro Pro Leu Ser Gln Arg His Thr
  180 185 190
- Phe Pro Glu Val Gln Arg Met Pro Ser Gln Ile Ser Leu Arg Asp Leu
  195 200 205
- Ser Glu Val Leu Glu Ala Glu Lys Val Pro His Asn Gln Arg Lys Pro 210 215 220
- Glu Ser Thr His Leu Leu Glu Asn Gln Asn Thr Gln Glu Ile Pro Leu 225 230 235 240
- Ala Ile Ser Ser Ser Ser Phe Thr Thr Ser Asn His Ser Val Gln Asn 245 250 255
- Arg Asp His Arg Gly Gly Met Gln Pro Cys Ser Pro Gln Arg Cys Gln 260 265 270
- Pro Pro Ala Ser Cys Ser Pro His Glu Asn Ile Leu Pro Tyr Lys Tyr 275 280 285
- Thr Ser Trp Arg Pro Pro Phe Pro Lys Arg Ser Asp Arg Lys Asp Val 290 295 300
- Gln His Asn Glu Trp Tyr Ile Gly Glu Tyr Ser Arg Gln Ala Val Glu 305 310 315 320
- Glu Ala Phe Met Lys Glu Asn Lys Asp Gly Ser Phe Leu Val Arg Asp 325 330 335
- Cys Ser Thr Lys Ser Lys Glu Glu Pro Tyr Val Leu Ala Val Phe Tyr

Glu Asn Lys Val Tyr Asn Val Lys Ile Arg Phe Leu Glu Arg Asn Gln 355 360 365

Gln Phe Ala Leu Gly Thr Gly Leu Arg Gly Asp Glu Lys Phe Asp Ser 370 375 380

Val Glu Asp Ile Ile Glu His Tyr Lys Asn Phe Pro Ile Ile Leu Ile 385 390 395 400

Asp Gly Lys Asp Lys Thr Gly Val His Arg Lys Gln Cys His Leu Thr 405 410 415

Gln Pro Leu Pro Leu Thr Arg His Leu Leu Pro Leu
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<210> 5

<211> 2540

<212> DNA

<213> HUMAN

<220>

<223> HUMAN MIST FULL-LENGTH CDNA SEQUENCE OF SPLICE VARIANT CLONE #12, NUCLEIC ACID SEQUENCE

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<210> 6

<211> 353

<212> PRT

<213> HUMAN

<220>

<223> HUMAN MIST SPLICE VARIANT CLONE #12, TRANSLATED AMINO ACID SEQUENCE

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Lys Glu Ser Glu Tyr Ala Asp Thr His Tyr Phe Lys Val Ala Met Asp
20 25 30

Thr Pro Leu Pro Leu Asp Thr Arg Thr Ser Ile Ser Ile Gly Gln Pro
35 40 45

Thr Trp Asn Thr Gln Thr Arg Leu Glu Arg Val Asp Lys Pro Ile Ser
50 55 60

Lys Asp Val Arg Ser Gln Asn Ile Lys Gly Asp Ala Ser Val Arg Lys

Asn	Lys	Ile	Pro	Leu	Pro	Pro	Pro	Arg	Pro	Leu	Ile	Thr	Leu	Pro	Lys
				85					90					95	

- Lys Tyr Gln Pro Leu Pro Pro Glu Pro Glu Ser Ser Arg Pro Pro Leu
  100 105 110
- Ser Gln Arg His Thr Phe Pro Glu Val Gln Arg Met Pro Ser Gln Ile 115 120 125
- Ser Leu Arg Asp Leu Ser Glu Val Leu Glu Ala Glu Lys Val Pro His 130 135 140
- Asn Gln Arg Lys Pro Glu Ser Thr His Leu Leu Glu Asn Gln Asn Thr 145 150 155 160
- Gln Glu Ile Pro Leu Ala Ile Ser Ser Ser Ser Phe Thr Thr Ser Asn 165 170 175
- His Ser Val Gln Asn Arg Asp His Arg Gly Gly Met Gln Pro Cys Ser 180 185 190
- Pro Gln Arg Cys Gln Pro Pro Ala Ser Cys Ser Pro His Glu Asn Ile 195 200 205
- Leu Pro Tyr Lys Tyr Thr Ser Trp Arg Pro Pro Phe Pro Lys Arg Ser 210 215 220
- Asp Arg Lys Asp Val Gln His Asn Glu Trp Tyr Ile Gly Glu Tyr Ser 225 230 235 240
- Arg Gln Ala Val Glu Glu Ala Phe Met Lys Glu Asn Lys Asp Gly Ser 245 250 255
- Phe Leu Val Arg Asp Cys Ser Thr Lys Ser Lys Glu Glu Pro Tyr Val 260 265 270
- Leu Ala Val Phe Tyr Glu Asn Lys Val Tyr Asn Val Lys Ile Arg Phe 275 280 285
- Leu Glu Arg Asn Gln Gln Phe Ala Leu Gly Thr Gly Leu Arg Gly Asp 290 295 300
- Glu Lys Phe Asp Ser Val Glu Asp Ile Ile Glu His Tyr Lys Asn Phe 305 310 315 320
- Pro Ile Ile Leu Ile Asp Gly Lys Asp Lys Thr Gly Val His Arg Lys

335

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<211> 8
<212> PRT
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<220>
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Leu

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<210> 7
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<213> HUMAN

<223> PHOSPHOPEPTIDE DERIVED FROM THE SEQUENCE OF HUMAN SLP-76

<221> MOD\_RES

<222> (3)

<223> PHOSPHORYLATION; TYR IN POSITION #3 IS PHOSPHORYLATED.

<400> 7

Asp Asp Tyr Glu Ser Pro Asn Asp

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<210> 8

<211> 21

<212> DNA

<213> HUMAN

<220>

<223> PRIMER PY474

<400> 8

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<210> 9

<211> 19

<212> DNA

<213> HUMAN

<220>

<223> PRIMER PY475

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Glu Leu
<210> 24
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His Tyr
<210> 25
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<212> PRT
<213> HUMAN
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Phe
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<212> PRT
<213> HUMAN
<400> 26
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                                     10
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 Glu Glu Thr Trp Gln Ser Ile Lys Ile Leu Pro Ala Arg
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 Ile Lys Gly Asp Ala Ser Val Arg Lys Asn Lys Ile Pro
                  5
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  <212> PRT
  <213> HUMAN
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<210> 30
<211> 13
<212> PRT
<213> HUMAN
<400> 30
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                                     10
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<400> 31
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<400> 32
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                  5
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<400> 34
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                                      10
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